

61 121 181 241 301 361 421	cgagaaaaggtgacgegggcegggaggeggeggeggegggeggeeeeeeee	
457	cggcggcgagcgggggccATGCAGGCGCGCTACTCCGTGTCCAGCCCCAACTCCMETGlnAlaArgTyrSerValSerSerProAsnSer	12
511	CTGGGAGTGGTGCCCTACCTCGGCGGCGAGCAGAGCTACTACCGCGCGCG	30
565	GCGGCGGCCGGGGCGGCTACACCGCCATGCCGGCCCCCATGAGCGTGTACTCG AlaAlaAlaGlyGlyTyrThrAlaMETProAlaProMETSerValTyrSer	48
619	CACCCTGCGCACGCCGAGCAGTACCCGGGCGGCATGGCCCGCGCCTACGGGCCC HisProAlaHisAlaGluGlnTyrProGlyGlyMETAlaArgAlaTyrGlyPro	66
673	TACACGCCGCAGCCGAGCCCAAGGACATGGTGAAGCCGCCCTATAGCTACATC TyrThrProGlnProGlnProLysAspMETValLysProProTyrSerTyrIle	84
727	GCGCTCATCACCATGGCCATCCAGAACGCCCCGGACAAGAAGATCACCCTGAAC AlaLeuIleThrMETAlaIleGlnAsnAlaProAspLysLysIleThrLeuAsn	102
781	GGCATCTACCAGTTCATCATGGACCGCTTCCCCTTCTACCGGGACAACAAGCAGGlyIleTyrGlnPheIleMETAspArgPheProPheTyrArgAspAsnLysGln	120
835	GGCTGGCAGAACAGCATCCGCCACAACCTCTCGCTCAACGAGTGCTTCGTCAAG GlyTrpGlnAsnSerIleArgHisAsnLeuSerLeuAsnGluCysPheValLvs	138
889	GTGCCGCGCGACGAGAGAGCCGGGCAAGGGCAGCTACTGGACGCTGGACCCG ValProArgAspAspLysLysProGlyLysGlySerTyrTrpThrLeuAspPro	15
943	GACTCCTACAACATGTTCGAGAACGGCAGCTTCCTGCGGCGGCGGCGCGCGC	174
997	AAGAAGAAGGACGCGGTGAAGGACAAGGAGGAGAAGGACAGGCTGCACCTCAAG LysLysAspAlaValLysAspLysGluGluLysAspArgLeuHisLeuLys	19:
1051	GAGCCGCCCCGCCCGGCCAGCCCAGCCCCGCCCGCCGCAGCA	21
1105	GGCAACGCGCCCGGTCCGCAGCCGCCCGTGCGCATCCAGGACATCAAGACC GlyAsnAlaProGlyProGlnProProProValArgIleGlnAspIleLysThr	22
1159	GAGAACGGTACGTGCCCTCGCCGCCCCAGCCCCTGTCCCCGGCCGCCGCCCTG GluAsnGlyThrCysProSerProProGlnProLeuSerProAlaAlaAlaLeu	24
1213	GGCAGCGGCAGCCGCCGCGGTGCCCAAGATCGAGAGCCCCGACAGCAGCAGC GlySerGlySerAlaAlaAlaValProLysIleGluSerProAspSerSerSer	26
1267	AGCAGCCTGTCCAGCGGGAGCAGCCCCCCGGGCAGCCTGCCGTCGGCGCGCGC	283
1321	CTCAGCCTGGACGGTGCGGATTCCGCGCCGCCGCCGCCCCCCCC	30
1375	CCGCCGCACCATAGCCAGGGCTTCAGCGTGGACAACATCATGACGTCGCTGCGG ProProHisHisSerGlnGlyPheSerValAspAsnIleMETThrSerLeuArg	31
1429	GGGTCGCCGCAGAGCGCGGCGCGGAGCTCAGCTCCGGCCTTCTGGCCTCGGCG GlySerProGlnSerAlaAlaAlaGluLeuSerSerGlyLeuLeuAlaSerAla	33
1483	GCCGCGTCCTCGCGCGCGGGGATCGCACCCCCGCTGGCGCTCGGCGCCTACTCG AlaAlaSerSerArgAlaGlyIleAlaProProLeuAlaLeuGlyAlaTyrSer	35
1537	CCCGGCCAGAGCTCCCTCTACAGCTCCCCTGCAGCCAGACCTCCAGCGGGC ProGlyGlnSerSerLeuTyrSerSerProCysSerGlnThrSerSerAlaGly	37

Fig. 1A

1591	AGCTCGGGCGGCGGCGGCGCGCGGGGGCCCGCGGGGGCGCGCCC SerSerGlyGlyGlyGlyGlyAlaGlyAlaAlaGlyGlyAlaGlyGlyAla	390
1645	GGGACCTACCACTGCAACCTGCAAGCCATGAGCCTGTACGCGGCCGGC	408
1699	GGGGGCCACTTGCAGGGCGCGCCCGGGGGGGGCGGCGGCTCGGCCGTGGACAAC GlyGlyHisLeuGlnGlyAlaProGlyGlyAlaGlyGlySerAlaValAspAsn	426
1753	CCCCTGCCCGACTACTCTCTGCCTCCGGTCACCAGCAGCAGCTCGTCGTCCCTG ProLeuProAspTyrSerLeuProProValThrSerSerSerSerSerLeu	444
1807	AGTCACGGCGGCGGCGGCGGCGGCGGGGGAGGCCAGGAGGCCGGCCACCA	462
1861	CCTGCGGCCCACCAAGGCCGCCTCACCTCGTGGTACCTGAACCAGGCGGGCG	480
1915	GACCTGGGCCACTTGGCAAGCGCGGCGGCGGCGGCGGCGGCGGCGCAGGCTACCCG AspLeuGlyHisLeuAlaSerAlaAlaAlaAlaAlaAlaAlaAlaGlyTyrPro	498
1969	GGCCAGCAGCAGAACTTCCACTCGGTGCGGGAGATGTTCGAGTCACAGAGGATCGlyGlnGlnAsnPheHisSerValArgGluMETPheGluSerGlnArgIle	516
2023	GGCTTGAACAACTCTCCAGTGAACGGGAATAGTAGCTGTCAAATGGCCTTCCCT GlyLeuAsnAsnSerProValAsnGlyAsnSerSerCysGlnMETAlaPhePro	534
2077	TCCAGCCAGTCTCTGTACCGCACGTCCGGAGCTTTCGTCTACGACTGTAGCAAG SerSerGlnSerLeuTyrArgThrSerGlyAlaPheValTyrAspCysSerLys	552
2131	TTTTGAcacaccctcaaagccgaactaaatcgaaccccaaagcaggaaaagcta PheSTP	554
223055555555555555555555555555555555555	aaggaacccatcaaggcaaaatcgaaactaaaaaaaaaa	こせせせ a a t a c a g g t g a t a t t a t c t a t c g

Fig. 1B

														, ·		•	
FKHR	FKHL10 FKHL13	FKHL3	FKHL4	FKHL6	FKHL5	FKHL17	FKHL8	FKHL9	FKHL15	FKHL12	FKHL11	FKHL18	FKHL14	Mutations	FKHL7	Forkhead	
WGNLSYADLITK.IESS.EKRLTLSQ.YEWMVKSVPYFKDKGDSNSSAKHSK.IR.QNEGTGKSSWWMLNPEGGKSGKSPRRAASMD	LMKLVRSAHGRLSQYVA.NNKS.ADKDKEDD TNPHATCM.ASKATSAKW.T.N.CYF.HADPTKIEKDEGF.RIQYAERLLS.AFKKLPFVHIH	G.YEKPFNMRQS.E.RLEKNYEHDKHYDDNMS.DDV.IG.TTGKLSTTSR	T. N. M. MEGGERT E. VN. V. E	F.GAYKVI.L.KGLGRH	F.GSYKVI.L.KGLGRH	DIEPGNN.	IQS.K.RLSE.CESGYEKFPADIEPGNN.	YRKFPDIEPGRN.	TDL.IEAGRN.	AARTEASPRKTDEPGNN.	DEQRVEKGREKGR	TTEPT		S M L	PKDMVKPPYSYIALITMAIQNAPDKKITLNGIYQFIMDRFPFYRDNKQGWQNSIRHNLSLNECFVKVPRDDKKPGKGSYWTLDPDSYNMFENGSFLRRRRRFKKKD	SYTHASN.TRMLSELQ.Q.RSF.DITPDFHLGCYQKCDK	Helix 1Helix 2Helix 3Wing 1Wing 2

Hig. 2

	Embryo			AA819240 AA964464	ლ ლ		Rat Rat	<i>ω</i>	UI-R-AO-al-b-03 UI-R-E1-go-e-12
1					سِ				mo83c06
	Embryo, 11.5 dy				<u>س</u>	psport1			vc85b07
<u>۔</u> ن	Embryo, 13.5-14.5 dy				<u>س</u>	pT7T3D	Mouse	403237	me94t07
い ロ ロ	×			AA739434	ũ	pr7r3D	Mouse	1226133	vv53d11
	bryo, 13.5-14.5			W91182	<u>.</u>	pt7t3d	Mouse	419796	mt72a07
	Embryo, 13.5-14.5 dy		W57082		<u>ω</u>	1	Mouse	372142	md53e12
	ummary Gland, 4	936	AA458089		ω	pr7r3D	Mouse	864300	vg45c07
	ummary Gland, 4		AA759405	AA960591	<u>ب</u>		Mouse	1248576	vw64c01
	Myotubes			AA673797	ທຸ		Mouse	1180061	vn08t03
	Kidney, 6 wk			AA276025	سِ	pT7T3D	Mouse	776052	vc30a07
l	Aorta			D57248	ယ္		Human		GEN-288A05
	NCI_CGAP_K1d3		AA886687		u		Human	1500703	oj3910 4
	Aorta			D56550	<u>س</u>		Human		GEN-206f07
	NCI_CGAP_PT22		AA688135		<u>س</u>		Human	1220412	nv16g07
	Placenta, 8 to 9 wk			N40582	ω -	pT7T3D	Human	258167	yw76d12
	NCI_CGAP_Pry			AA551599	<u>ო</u>		Human	996558	nj57a04
	Fetal Cochlea	475	N22552	N75774		pBlue SK-	Human	253733	yw30d03
	Embryo, 9 wk	: !		AA334694	<u>س</u>		Human		EST38957
	retal Heart			AA348051	س -		Human		EST54452
	Fetal Cochlea			н89575		pBlue SK-	Human	253556	yw28c11
	Placenta, 8 to 9 wk		N25867		<u>ب</u>		Human	258335	yw78b12
	•		AA902429		<u>س</u>		Human	1521276	ok90g07
	etal Heart			W94714	ω		Human	358885	ze13t07
	Fetal Heart, 19 wk	919	AA022755	AA022618	<u>ب</u>	pr7r3D	Human	364392	ze71a01
	etal Heart, 19		W73917		ω <u></u>		Human	346079	zd71b12
	NCI_CGAP_GC4		AA865139		uٍ		Human	1469849	oh48b09
	Wilms Tumor		AA776534		سِ		Human	1156628	ah14c11
	Fetal Heart, 19 wk			W77980	<u>س</u>		Human	346077	zd71b11
	NCI_CGAP_Lu5		AA885880		<u>س</u>	ı	Human	1500423	oj36t08
	NhHMPu	722		AA495846	ယ္	pT7T3D	Human	768370	zw05a06
	Placenta, 8 to 9 wk		N25875		ယ္		Human	258359	vw78d12
	Fetal Heart		W94629		ယ္	•	Human	358885	ze13t07
	Placenta, 8 to 9 wk			N40575	ယ္	pT7T3D	Human	258143	yw76b12
	NhemPu		AA424466	AA424381	ω		Human	767110	zv90g12
	NhHMPu		AA424787		ယ္		Human	768274	zw04a06
	NhHMPu		AA232201	AA232742	u		Human	666326	zr45a08
	Arana contra	27.70	3" Sequence	5. Sequence	00 FOC	Vector	organism	Number	Name
:. 3		Insert			•		-	Image	Clone